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Is 26	1	MRHMQSVTRASSI	CGSGVKQVI	GKGHPHARV	VGARAKA	QIPEREE
Is 27	1	MRHMQSATRATL	VCGSGVKQI	IAKGHPNTR	VFGARAKA	QIPEREV
Is 26	45	LSVKPKMVRNTH	LNLPQERQAF	YRLL	LENEQIQEFL	SMDSCLRI
Is 27	45	LAAKPKITRITH	LNLPQERQAF	YRLL	LENELIQEFL	SMDSCLKI
Is 26	89	SDKYLIAMVLA	YFKRAAGLYT	SEYTTMN	FFVALYL	ANDMEEDEE
Is 27	89	SDKYLIAMVLA	YFKRA - GLYT	SEYTTMN	FFVALYL	ANDMEEDEE
Is 26	133	DYKYEI	FPWALGDSWREL	FPQFLRL	RDDFWAKMNY	RAVVSRRCC
Is 27	132	DYKYEI	FPWALGDSWREF	FPQFLRL	RDDFWAKMNY	RAVVSRRCC
Is 26	177	DEVMSKDPTH	HWAWLRDRP	MHHSGAMRGYL	RNEDDFF	PRGPGGLTP
Is 27	176	DEVMAKDP	THWAWLRDRPI	HHSGALRGYL	RNEDDFF	PRGPGGLTP
Is	221	ASCTLCHKAGV	CDSGGVSHNN	SSPEQEI	FHYTNREWSQ	ELLML
Is 27	220	ASCALCHKASV	CDSGGVSHDNSS -	PEQEI	FHYTNREWSQ	ELLIL
Is 26	265	PPELLLDPECT	HDLHI	LQEPLVGL	EPDGTAL	EWHL
Is 27	263	PPELLLDPESTY	DIHI	FQEPLVGL	EPDGAAL	EWHL

Fig.1

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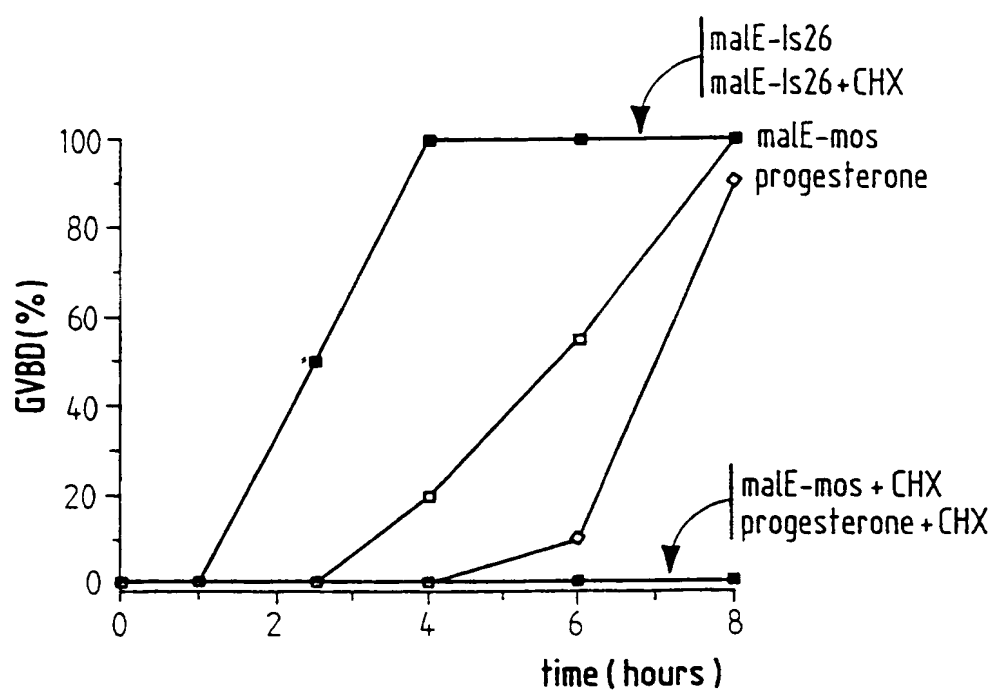


Fig. 2

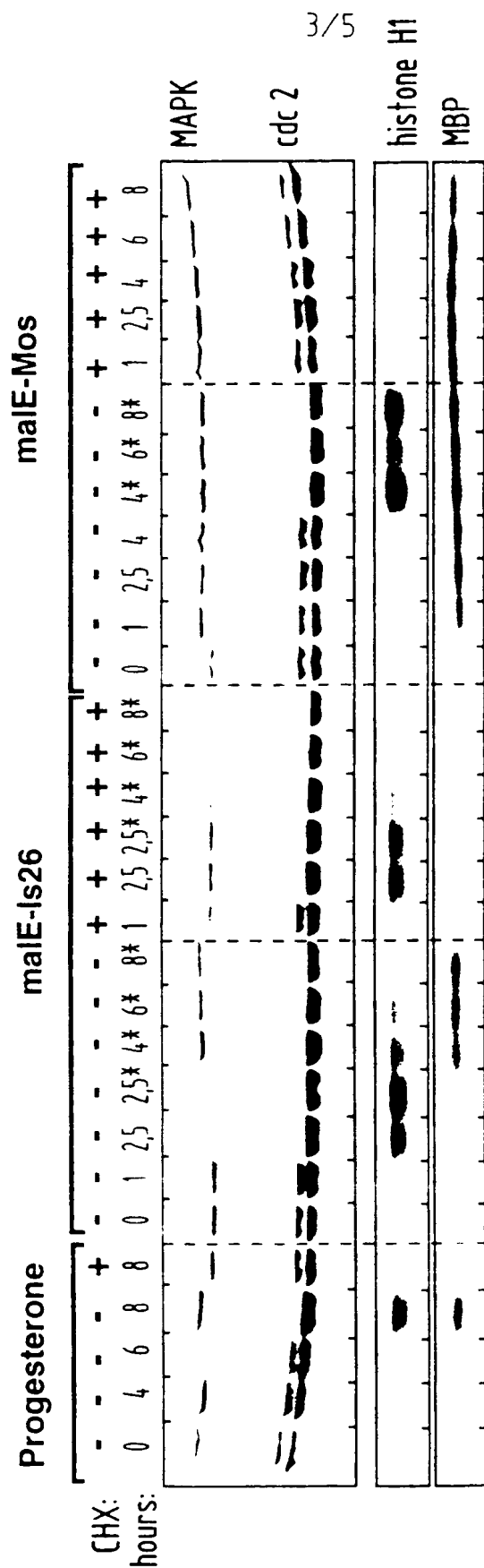


Fig. 3

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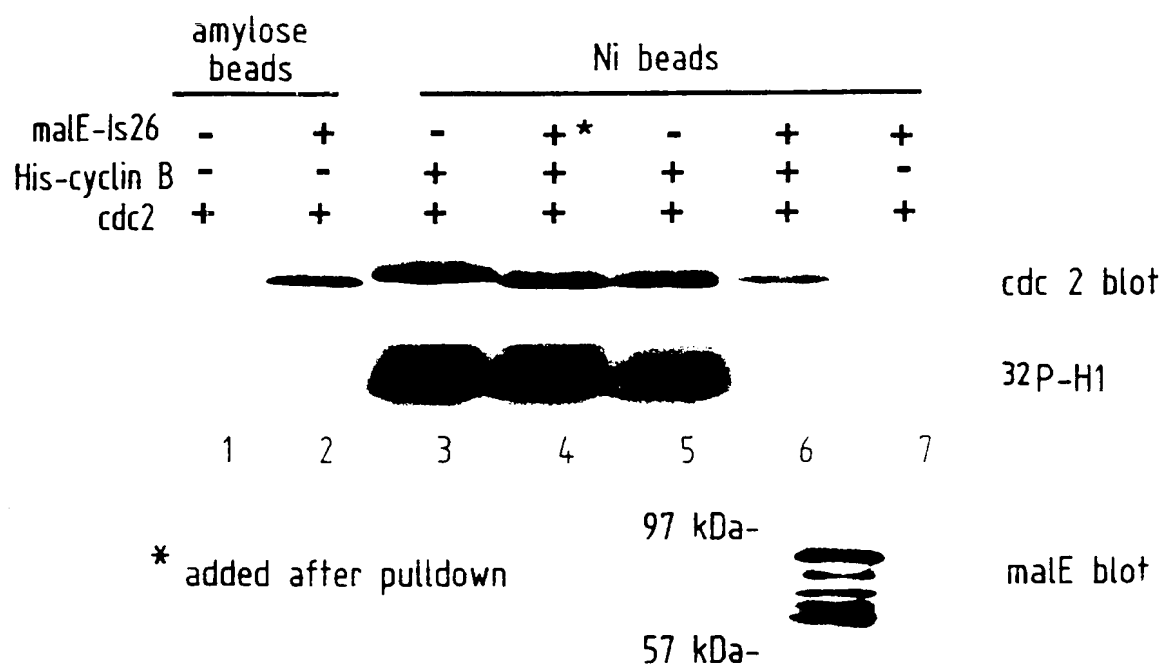


Fig. 4

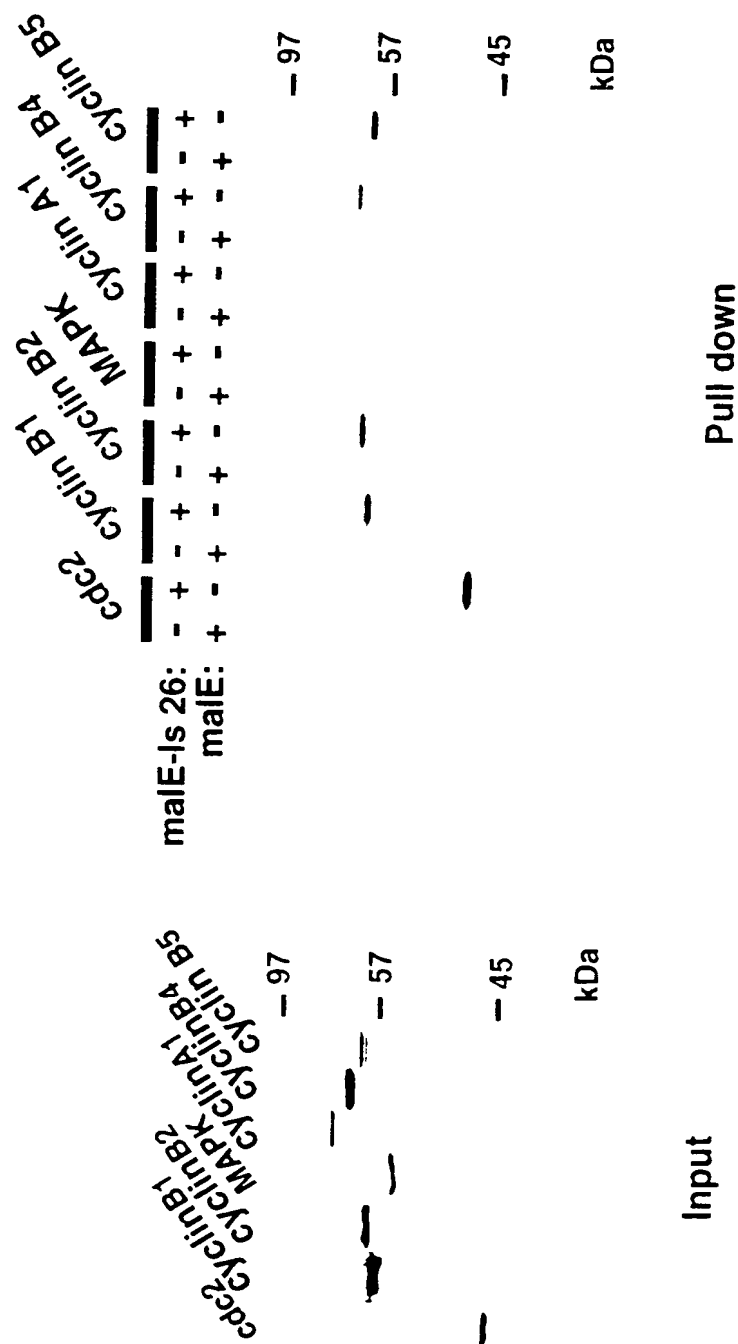


Fig. 5